STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/550,608
Source:	IFWP
Date Processed by STIC:	11/14/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

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- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm , EFS Submission User Manual ePAVE)
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Revised 01/10/06



IFWP

RAW SEQUENCE LISTING

DATE: 11/14/2006

PATENT APPLICATION: US/10/550,608

TIME: 08:51:21

Input Set : N:\RJAVED\10550608.txt

Output Set: N:\CRF4\11142006\J550608.raw

4 <110> APPLICANT: Progenika Biopharma, S.A.

6 <120> TITLE OF INVENTION: In vitro method to detect bladder transitional cell carcinoma

C--> 8 <140> CURRENT APPLICATION NUMBER: US/10/550,608

C--> 8 <141> CURRENT FILING DATE: 2005-09-26

0 <130> FILE REFERENCE:

8 <160> NUMBER OF SEQ ID NOS: 18

10 <170> SOFTWARE: PatentIn version 3.1

Does Ry - oly Corrected Lukefle Needed Cpg-1)

ERRORED SEQUENCES

282 <210> SEQ ID NO: 18

283 <211> LENGTH: 25

284 <212> TYPE: DNA

285 <213> ORGANISM: Artificial sequence

not Greet any to W--> 287 <220> FEATURE: synthetic DNA 288 <223> OTHER INFORMATION: probe sequence of the 31805 at probe set of Affymetrix, the

position of said probe in the mRNA sequence of the fgfr3 gene being

290 3720

293 <400> SEQUENCE: 18

294 gtggccagag gtgtcaccca aaccg

E--> 299/1

Pls delete

This lype of Error Occur subsequently Occur subsequently throughout the Sequence.

VERIFICATION SUMMARYDATE: 11/14/2006PATENT APPLICATION: US/10/550,608TIME: 08:51:22

Input Set: N:\RJAVED\10550608.txt
Output Set: N:\CRF4\11142006\J550608.raw

L:8 M:270 C: Current Application Number differs, Replaced Current Application No L:8 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:0 M:201 W: Mandatory field data missing, <130> FILE REFERENCE L:17 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:32 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:47 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:63 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:79 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:95 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:111 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:127 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:143 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:159 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:175 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:191 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:207 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:223 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:239 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:255 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:271 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:287 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:299 M:254 E: No. of Bases conflict, this line has no nucleotides.